

10/566679

SEQUENCE LISTING

AP20 Rec'd PCT/PTO 01 FEB 2006

<110> CENTRE NATIONAL DE LA RECHERCHE SCIENTIFIQUE

<120> NOVEL ANTI-ANGIOGENIC AGENT AND ITS USE, IN PARTICULAR
WITHIN THE FRAMEWORK OF THE TREATMENT OF CANCER

<130> WOB 03 AW CNR GIOG

<150> FR 03/09506

<151> 2004-08-01

<160> 12

<170> PatentIn version 3.1

<210> 1

<211> 2389

<212> DNA

<213> homo sapiens

<220>

<221> CDS

<222> (73)..(1143)

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Met Gln Ser Val Gln Ser Thr Ser Phe Cys Leu Arg Lys

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cag tgc ctt tgc ctg acc ttc ctg ctt ctc cat ctc ctg gga cag gtc 159

Gln Cys Leu Cys Leu Thr Phe Leu Leu Leu His Leu Leu Gly Gln Val

15 20 25

gct gcg act cag cgc tgc cct ccc cag tgc ccg gcc cgg tgc cct gcg 207

Ala Ala Thr Gln Arg Cys Pro Pro Gln Cys Pro Gly Arg Cys Pro Ala

30 35 40 45

acg ccg ccg acc tgc gcc ccc ggg gtg cgc gcg gtg ctg gac gcc tgc 255

Thr Pro Pro Thr Cys Ala Pro Gly Val Arg Ala Val Leu Asp Gly Cys

50 55 60

tca tgc tgt ctg gtg tgt gcc cgc cag cgt gcc gag agc tgc tca gat 303

Ser Cys Cys Leu Val Cys Ala Arg Gln Arg Gly Glu Ser Cys Ser Asp

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ctg gag cca tgc gac gag agc agt gcc ctc tac tgt gat cgc agc gcg 351

Leu Glu Pro Cys Asp Glu Ser Ser Gly Leu Tyr Cys Asp Arg Ser Ala

80 85 90

gac ccc agc aac cag act gcc atc tgc acg gcg gta gag gga gat aac 399

Asp Pro Ser Asn Gln Thr Gly Ile Cys Thr Ala Val Glu Gly Asp Asn

95 100 105

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Cys Val Phe Asp Gly Val Ile Tyr Arg Ser Gly Glu Lys Phe Gln Pro

110 115 120 125

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cca aga aaa gtt gag gtg cct gga gag tgc tgt gaa aag tgg atc tgt Pro Arg Lys Val Glu Val Pro Gly Glu Cys Cys Glu Lys Trp Ile Cys 160 165 170	591
ggc cca gat gag gag gat tca ctg gga ggc ctt acc ctt gca gct tac Gly Pro Asp Glu Glu Asp Ser Leu Gly Gly Leu Thr Leu Ala Ala Tyr 175 180 185	639
agg cca gaa gcc acc cta gga gta gaa gtc tct gac tca agt gtc aac Arg Pro Glu Ala Thr Leu Gly Val Glu Val Ser Asp Ser Ser Val Asn 190 195 200 205	687
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Gln Arg Cys Pro Pro Gln Cys Pro Gly Arg Cys Pro Ala Thr Pro Pro
35 40 45
Thr Cys Ala Pro Gly Val Arg Ala Val Leu Asp Gly Cys Ser Cys Cys
50 55 60
Leu Val Cys Ala Arg Gln Arg Gly Glu Ser Cys Ser Asp Leu Glu Pro
65 70 75 80
Cys Asp Glu Ser Ser Gly Leu Tyr Cys Asp Arg Ser Ala Asp Pro Ser
85 90 95
Asn Gln Thr Gly Ile Cys Thr Ala Val Glu Gly Asp Asn Cys Val Phe
100 105 110
Asp Gly Val Ile Tyr Arg Ser Gly Glu Lys Phe Gln Pro Ser Cys Lys
115 120 125
Phe Gln Cys Thr Cys Arg Asp Gly Gln Ile Gly Cys Val Pro Arg Cys
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Gln Leu Asp Val Leu Leu Pro Glu Pro Asn Cys Pro Ala Pro Arg Lys
145 150 155 160
Val Glu Val Pro Gly Glu Cys Cys Glu Lys Trp Ile Cys Gly Pro Asp
165 170 175

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Glu Glu Asp Ser Leu Gly Gly Leu Thr Leu Ala Ala Tyr Arg Pro Glu
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 Ala Thr Leu Gly Val Glu Val Ser Asp Ser Ser Val Asn Cys Ile Glu
 195 200 205
 Gln Thr Thr Glu Trp Thr Ala Cys Ser Lys Ser Cys Gly Met Gly Phe
 210 215 220
 Ser Thr Arg Val Thr Asn Arg Asn Arg Gln Cys Glu Met Leu Lys Gln
 225 230 235 240
 Thr Arg Leu Cys Met Val Arg Pro Cys Glu Gln Glu Pro Glu Gln Pro
 245 250 255
 Thr Asp Lys Lys Gly Lys Lys Cys Leu Arg Thr Lys Lys Ser Leu Lys
 260 265 270
 Ala Ile His Leu Gln Phe Lys Asn Cys Thr Ser Leu His Thr Tyr Lys
 275 280 285
 Pro Arg Phe Cys Gly Val Cys Ser Asp Gly Arg Cys Cys Thr Pro His
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 Asn Thr Lys Thr Ile Gln Ala Glu Phe Gln Cys Ser Pro Gly Gln Ile
 305 310 315 320
 Val Lys Lys Pro Val Met Val Ile Gly Thr Cys Thr Cys His Thr Asn
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 acc tgc gcc ccc ggg gtg cgc gcg gtg ctg gac ggc tgc tca tgc tgt 96
 Thr Cys Ala Pro Gly Val Arg Ala Val Leu Asp Gly Cys Ser Cys Cys
 20 25 30

tgc	gac	gag	agc	agt	ggc	ctc	tac	tgt	gat	cgc	agc	gcg	gac	ccc	agc	192
Cys	Asp	Glu	Ser	Ser	Gly	Leu	Tyr	Cys	Asp	Arg	Ser	Ala	Asp	Pro	Ser	
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Leu Val Cys Ala Arg Gln Arg Gly Glu Ser Cys Ser Asp Leu Glu Pro
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Cys Asp Glu Ser Ser Gly Leu Tyr Cys Asp Arg Ser Ala Asp Pro Ser
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Asn Gln Thr Gly Ile Cys Thr Ala
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cag cca agc tgc aaa ttc cag tgc acc tgc aga gat ggg cag att ggc 96
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                20            25          30  
  
Cys Val Pro Arg Cys Gln Leu Asp Val Leu Leu Pro Glu Pro Asn Cys  
        35              40          45  
  
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1				5					10						15				
atg ggg ttc tcc acc cgg gtc acc aat agg aac cgt caa tgt gag atg																			96
Met Gly Phe Ser Thr Arg Val Thr Asn Arg Asn Arg Gln Cys Glu Met																			
			20					25						30					

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1          5          10          15
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aac tgc acc agc ctg cac acc tac aag ccc agg ttc tgt ggg gtc tgc 96
Asn Cys Thr Ser Leu His Thr Tyr Lys Pro Arg Phe Cys Gly Val Cys
20 25 30

agt gat ggc cgc tgc tgc act ccc cac aat acc aaa acc atc cag gca 144
 Ser Asp Gly Arg Cys Cys Thr Pro His Asn Thr Lys Thr Ile Gln Ala
 35 40 45

gag ttt cag tgc tcc cca ggg caa ata gtc aag aag cca gtg atg gtc 192
Glu Phe Gln Cys Ser Pro Gly Gln Ile Val Lys Lys Pro Val Met Val
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 Ile Gly Thr Cys Thr Cys His Thr Asn Cys Pro
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 Val Asn Cys Ile Glu Gln Thr Thr Glu Trp Thr Ala Cys Ser Lys Ser
 20 25 30
 tgt ggt atg ggg ttc tcc acc cgg gtc acc aat agg aac cgt caa tgt 144
 Cys Gly Met Gly Phe Ser Thr Arg Val Thr Asn Arg Asn Arg Gln Cys
 35 40 45
 gag atg ctg aaa cag act cgg ctc tgc atg gtg cgg ccc tgt gaa caa 192
 Glu Met Leu Lys Gln Thr Arg Leu Cys Met Val Arg Pro Cys Glu Gln
 50 55 60
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 Glu Pro Glu Gln Pro Thr Asp Lys Lys Gly Lys Lys Cys Leu Arg Thr
 65 70 75 80
 aag aag tca ctc aaa gcc atc cac ctg cag ttc aag aac tgc acc agc 288
 Lys Lys Ser Leu Lys Ala Ile His Leu Gln Phe Lys Asn Cys Thr Ser
 85 90 95

ctg cac acc tac aag ccc agg ttc tgt ggg gtc tgc agt gat ggc cgc 336
 Leu His Thr Tyr Lys Pro Arg Phe Cys Gly Val Cys Ser Asp Gly Arg
 100 105 110

tgc tgc act ccc cac aat acc aaa acc atc cag gca gag ttt cag tgc 384
 Cys Cys Thr Pro His Asn Thr Lys Thr Ile Gln Ala Glu Phe Gln Cys
 115 120 125

tcc cca ggg caa ata gtc aag aag cca gtg atg gtc att ggg acc tgc 432
 Ser Pro Gly Gln Ile Val Lys Lys Pro Val Met Val Ile Gly Thr Cys
 130 135 140

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 Thr Cys His Thr Asn Cys Pro Lys Asn Asn Glu Ala Phe Leu Gln Glu
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<210> 12
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 <213> Homo sapiens

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 20 25 30

Cys Gly Met Gly Phe Ser Thr Arg Val Thr Asn Arg Asn Arg Gln Cys
 35 40 45

Glu Met Leu Lys Gln Thr Arg Leu Cys Met Val Arg Pro Cys Glu Gln
 50 55 60

Glu Pro Glu Gln Pro Thr Asp Lys Lys Gly Lys Lys Cys Leu Arg Thr
 65 70 75 80

Lys Lys Ser Leu Lys Ala Ile His Leu Gln Phe Lys Asn Cys Thr Ser
 85 90 95

Leu His Thr Tyr Lys Pro Arg Phe Cys Gly Val Cys Ser Asp Gly Arg
 100 105 110

Cys Cys Thr Pro His Asn Thr Lys Thr Ile Gln Ala Glu Phe Gln Cys
 115 120 125

Ser Pro Gly Gln Ile Val Lys Lys Pro Val Met Val Ile Gly Thr Cys
 130 135 140

Thr Cys His Thr Asn Cys Pro Lys Asn Asn Glu Ala Phe Leu Gln Glu
 145 150 155 160

Leu Glu Leu Lys Thr Thr Arg Gly Lys Met
 165 170